

4/25

		A	
ZNFN3A1 : 1	MEPLKVEKFATANRGNGLRAVTPLRPGELLFRSDPLAYTVCKGSRGVVCDRCLLGKEKLM	60	
	ME LKVEKF TANRGNGLRAV PLRPGELLFRSDPLAYTVCKGSRGVVCDRCLLGKEKLM		
AK010447: 1	MEALKVEKFTTANRGNGLRAVAPLRPGELLFRSDPLAYTVCKGSRGVVCDRCLLGKEKLM	60	
ZNFN3A1 : 61	RCSQCRVAKYCSAKCQKKAWPDHKRECKCLKSCKPRYPDSVRLLGRVVKLMDGAPSES	120	
	RCSQCR+AKYCSAKCQKKAWPDH+REC CLKSCKPRYPDSVRLLGRV+ KLMD PSES		B
AK010447: 61	RCSQCRVAKYCSAKCQKKAWPDHRECSCLKSCKPRYPDSVRLLGRVIVKLMDEKPSES	120	
ZNFN3A1 : 121	EKLYSFYDLESNINKLTEDKKEGLRQLVMTFQHFMRREETODASQLPPAFDLFEAFKVIC	180	
	EKLYSFYDLESNI+KLTEDKKEGLRQLVMTFQHFMRREETODASQLPP+FDLFEAFKVIC		
AK010447: 121	EKLYSFYDLESNISKLTEDKKEGLRQLVMTFQHFMRREETODASQLPPSFDLFEAFKVIC	180	
ZNFN3A1 : 181	NSFTICNAEMQEVGVGLYPSISLNHSCDPNCSTVFNGPHLLLRAVRDIEVGEELTICYL	240	
	NSFTICNAEMQEVGVGLYPS+SLLNHSCDPNCSTVFNGPHLLLRAVR+IE GEELTICYL		
AK010447: 181	NSFTICNAEMQEVGVGLYPSISLNHSCDPNCSTVFNGPHLLLRAVREIEAGEELTICYL	240	
ZNFN3A1 : 241	DMLMTSEERRKQLRDQYCFE CDCFRCTQDKDADMLTGDEQVWKEVQESLKKIEELKAHW	300	
	DMLMTSEERRKQLRDQYCFE CDC RCQTQDKDADMLTGDEQ+WKEVQESLKKIEELKAHW		
AK010447: 241	DMLMTSEERRKQLRDQYCFE CDCIRCQTQDKDADMLTGDEQIWKEVQESLKKIEELKAHW	300	
ZNFN3A1 : 301	KWEQVLAMCQAISSNSERLPDINIYQLKVLD CAMDACINLG LLEEALFYGTRTMEPYRI	360	
	KWEQVLA+CQAI+SNS RLPDINIYQLKVLD CAMDACINLG+LEEALFY RTMEPYRI		
AK010447: 301	KWEQVLALCQAIINSNSNRLPDINIYQLKVLD CAMDACINLG MLEEALFYAMRTMEPYRI	360	
ZNFN3A1 : 361	FFPGSHPVRGVQVMKVGLQLHQGMFPQAMKNLRLAFDIMRVTHGREHSLIEDLILLEE	420	
	FFPGSHPVRGVQVMKVGLQLHQGMFPQAMKNLRLAFDIM+VTHGREHSLIEDLILLEE		
AK010447: 361	FFPGSHPVRGVQVMKVGLQLHQGMFPQAMKNLRLAFDIMKVTHGREHSLIEDLILLEE	420	
ZNFN3A1 : 421	CDANIRAS	428	(SEQ ID NO: 1)
	CDANIRAS		(SEQ ID NO: 81)
AK010447: 421	CDANIRAS	428	(SEQ ID NO: 82)

FIG.2B

10/25

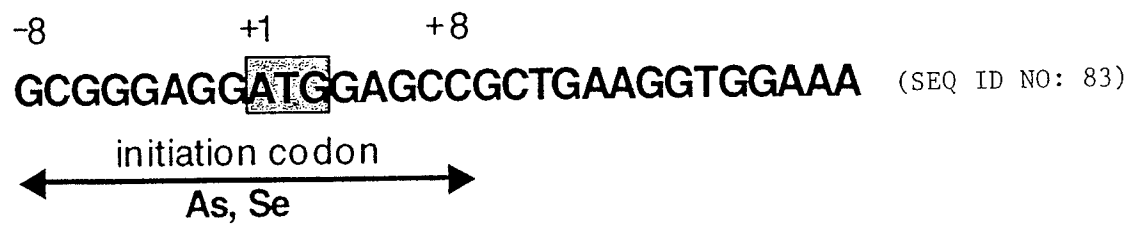


FIG.6A

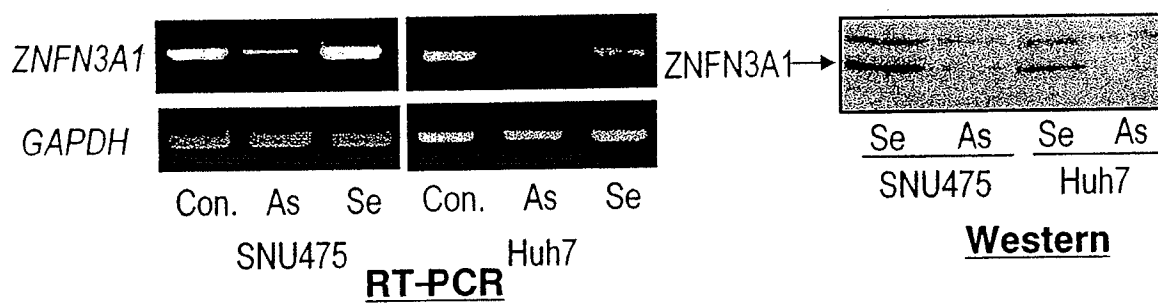


FIG.6B

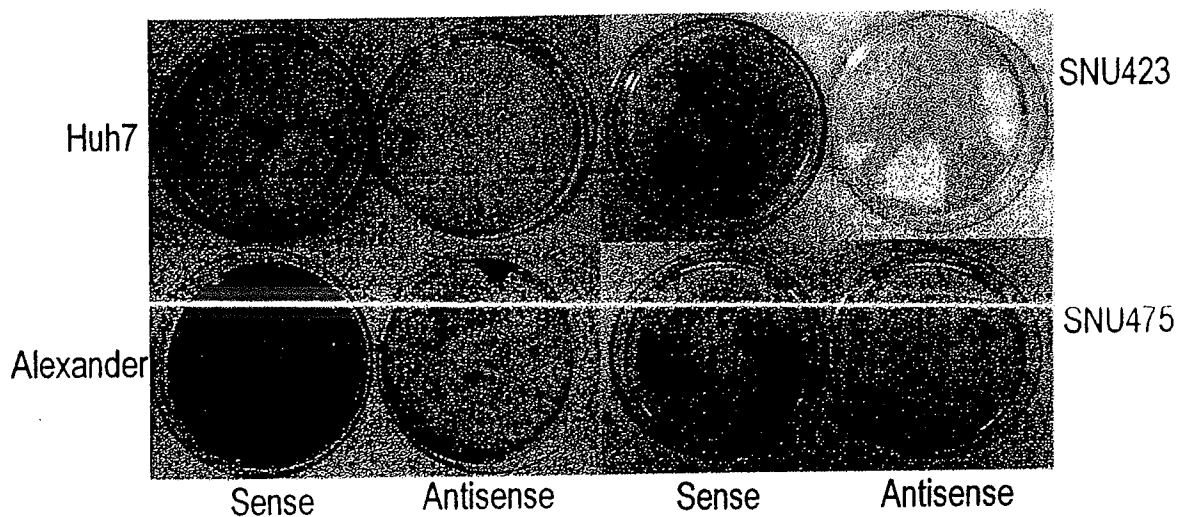


FIG.6C